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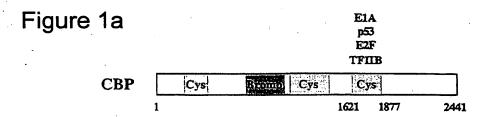
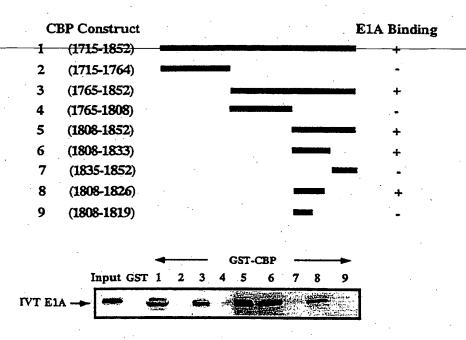


Figure 1b

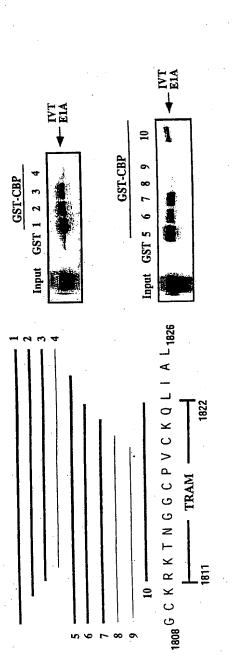


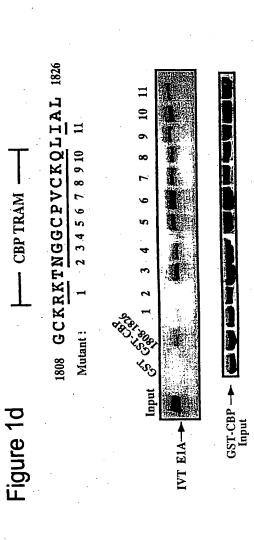
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Figure 1c





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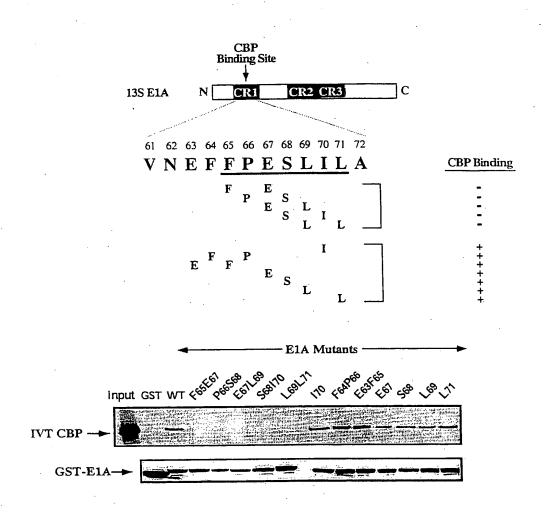


Figure 2a

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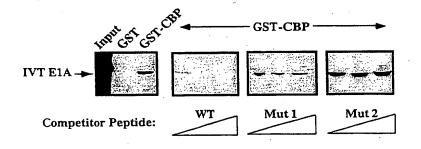
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#### Figure 2b

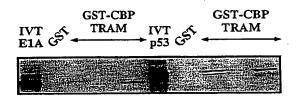
 WT
 :
 V N E F F P E S L I L

 Peptides:
 Mut 1 :
 V N E F F P A S A I L

 Mut 2 :
 V N E F A P A S A I A



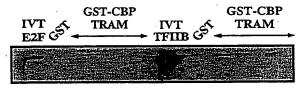
#### Figure 2c



E1A Competitor:

- WT Mut

WT Mut



E1A Competitor:

- WT Mut

WT Mut

## Figure 2d

TRAM F1A VNEFFPESLILAA
Interaction p53 SQETFSDLWKLLP
Motif E2F FDCDFGDLTPLDF

Consensus:

 $FX\frac{E}{D}XXXL$ 

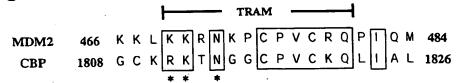


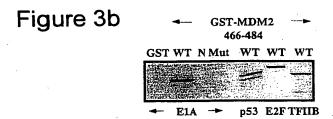
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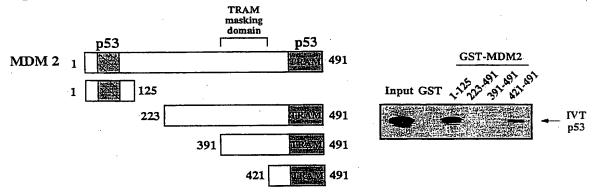




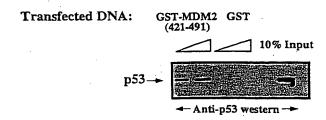




## Figure 3c



#### Figure 3d

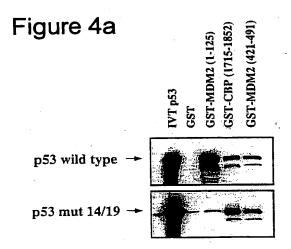


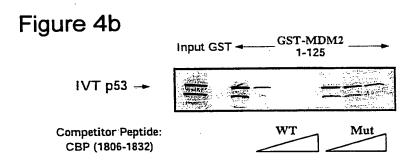
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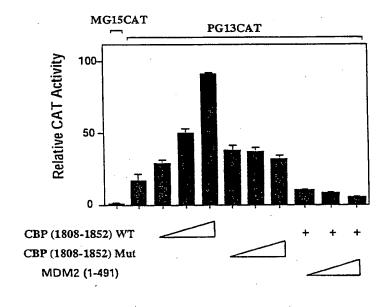
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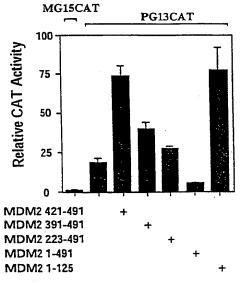




## Figure 4c



## Figure 4d



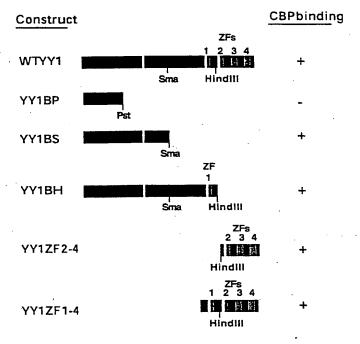


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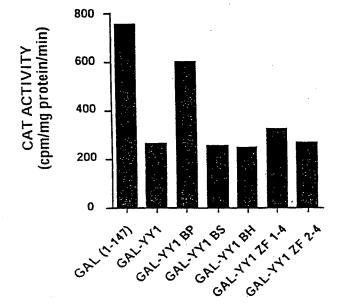
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## Figure 5b

## Figure 5c





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## Figure 6a

		E1A	VNEF <b>FPE</b> SLI <b>L</b> AA
		<b>p53</b>	SQET <b>FSDLWKL</b> LP
TRIMs	_	E2F	FDCD <b>FGDLTPL</b> DF
	ļ	TFIIB	MMNAFKEITTMAD
		YY1	AEDG <b>FED</b> OIL <b>I</b> PV
	_	YY1	CTKM <u>FRDNSAM</u> RK
		YY1	CGKA <b>FVE</b> SSK <b>L</b> KR
	•	MyoD	TTDD <b>FYD</b> DPC <b>F</b> DS

# Figure 6b

--- CBP TRAM ----

1808 GCKRKT NGGCPVCKQLIAL 1826 Mutant: 1 2 3 4 5 6 7 8 9 10 11

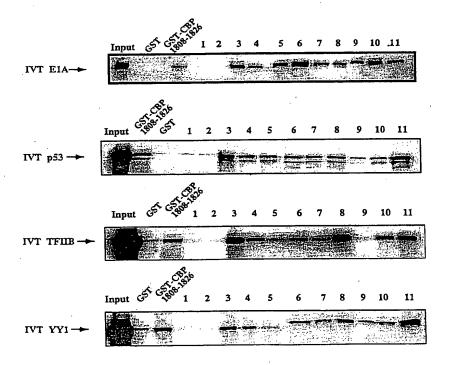






Figure 7a

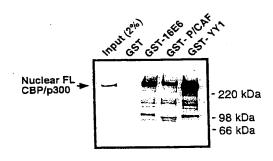
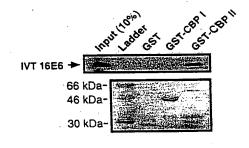
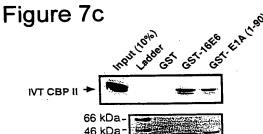
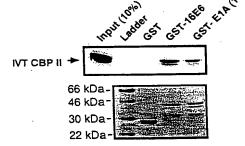


Figure 7b







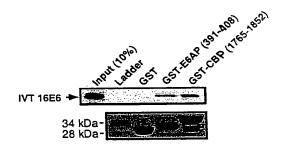


Figure 7d



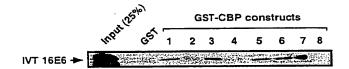
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	CBP constructs	E6 binding
1	1715 1870	+
2	1715 1764	-
3	1765 1852	+
4	1765 1808	-
5	1808 1852	+
6	1808 1833	+
7	1808 1826	+
8	1808 1819	-



CBP (1808-1826): GCKRKTNGGCPVCKQLIAL p300 (1770-1788): GCKRKTNGGCPICKQLIAL

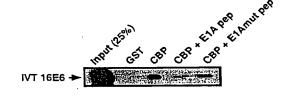
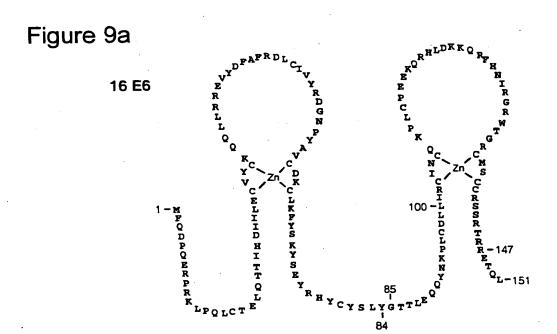


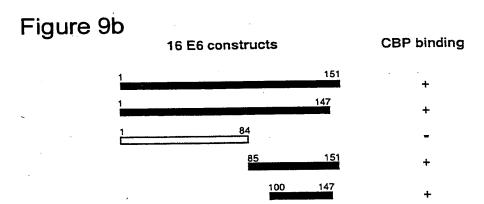
Figure 8

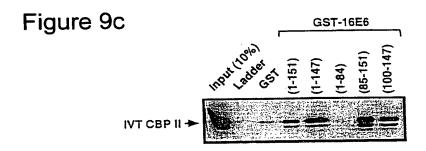


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Figure 10a

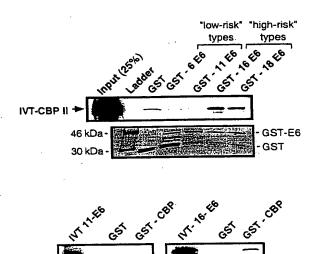
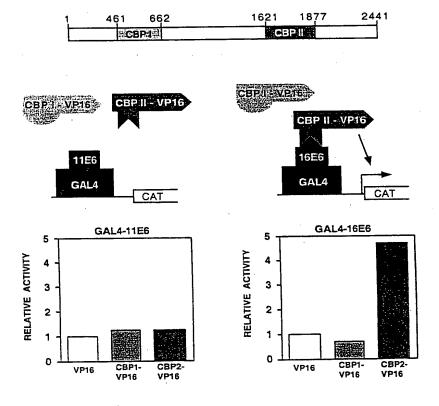


Figure 10b



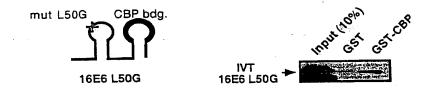
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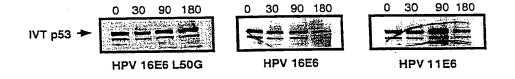


# Figure 11a





## Figure 11c



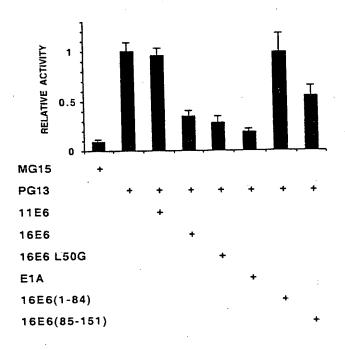


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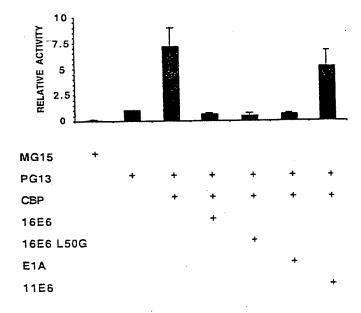


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Figure 12a



# Figure 12b





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TTC GAA CTA ACT AAT GGA a 5' HindIII stop (x3) [core HindIII] CCC GGG CTG CAG AAG CT<u>I GA</u>T TGA Tta gct GGG CCC GAC GTC TTC GAA CTA ACT AAT CTT Smal Pstl Hindti L E CTC GAG (GAG CTC (Xhol G S V D
gGA TCC GTC GAC C
cct agG CAG CTG G
BamHl Sal1

Figure 13